

Figure 1

GGCAGAGCTCTCCTCGTCCCTCCCTTCTCCACTGCAGCCTTTCTCTTAGCCCGAACCA 60
CTTCCTTCTTCTGCTTGTTCCTCCCTAGGGCGCGGAAGCTGAGTGCAGGGTTCAGACCCA 120
CGCGGCGAGCAGCTCTTCAGTGAAGAAGGAAGCAATCGGAGGGTCAGCAATGAACGTGGA 180
M N V E
GCATGAGGTTAACCTCCTGCTGGAGGAAATTCATCGTCTGGGTTCCAAAAATGCCGATGG 240
H E V N L L V E E I H R L G S K N A D G
GAAACTGAGTGTGAAGTTTGGGGTCCTCTTCCAAGACGACAGATGTGCCAATCTCTTTGA 300
K L S V K F G V L F Q D D R C A N L F E
AGCGTTGGTGGGAACCTCTGAAAGCCGCAAAACGAAGGAAGATTGTTACGTACGCAGGAGA 360
A L V G T L K A A K R R K I V T Y A G E
GCTGCTTTTGAAGGTGTTTCATGATGATGTTGACATTGTATTGCTGCAAGATTAATGTGG 420
L L L Q G V H D D V D I V L L Q D
TTTGCAGATCTGGGGGTATCTGGTAAACTGGAATAATTAAGTTAAAGGACAAACATGAAG 480
TTCCTTATGTATTTTATAGACCTTTGTAAACAAAAGGGGACTTGTTGAGAAGTCCTGTT 540
TTTATACCTTGGAGCAAAACATTACAATGTAAAAATAAACAAAACCTGTTATTTTTTTTT 600
TCTTAAGAAGGTAATCGGGAGACGTAGGCAATAAAATGTTTTTCAGAGGTGCGAAAAAGCT 660
TTTGTTTTCTTAAACCATTTCTTAGTCTCTGCCACACTTGACACTCCGTCAAAGTGAGAAG 720
CGAACTAAAGACCAACTGCGGTGGAAAATATTATGTTTATGTAATAAAAAAAAATCATGT 780

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Figure 2

GGCACGAGGCTTGAGCGCAGAAACACTTACTTTTCCCCCTACCCTGCTCCTCCTCCA 60
CAGCCGTCTTTCTCTTTGCCCTCAGCCACTTCCTTCCTTCGCCTCACCCCTCCCAGTGCAC 120
TGAAGAAGGTAACCGGGTCCAGACCCACGCGGCGCCAGTTCTCCGGCGGGAAGGAAAACC 180
GCGCAGAGAGGCAGCAATGAATGTGGATCACGAGGTTAACCTCTTAGTGGAGGAAATCA 240
M N V D H E V N L L V E E I H
TCGTTTGGGTTCAAAAAATGCTGATGGAAAAGTTAAGCGTGAAATTTGGGGTCCTCTCCG 300
R L G S K N A D G K L S V K F G V L F R
TGATGATAAATGTGCCAACCTCTTTGAAGCATTGGTAGGAACCTTTAAAGCTGCAAAACG 360
D D K C A N L F E A L V G T L K A A K R
AAGGAAGATTGTAACATATCCAGGAGAGCTGCTTCTGCAAGGTGTTTCATGATGATGTTGA 420
R K I V T Y P G E L L L Q G V H D D V D
CATTATATTACTGCAAGATTAATGTGGTTTACATATCTTTATGTACTGCCATTTTTTGGT 480
I I L L Q D
TCTGGTAAACTGGAATATAAAGTGAAAGAACAACATTTGAACATACTTAATGTATTTTTT 540
ATAGAACTTTGTAAACGAAAGGAGATTTCATGTTTTAGAAAGTCTGTCCTTTTTTATATCTT 600
GAAAGAAAATCTATGTATGATGCTATAAAATAAATCCTATTATTTTTTCTCAGGAATCTGG 660
TTAGGAATTGCAGGCAATGAGATTTTTTTCGGGGGCAGGGATGGGAATGTTTGTTCATAAA 720
TAATTAGACATTTTCTATAGATATTTGACATTCTGCGAAAGCAACAAGCAAACCTGAAGAC 780
CAACTCCTATGAGAAATATTATGATGTTTATGTAATAAAGACATGTAACCTGTCTT 835

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RatPSGen-13	-----GGCACGAGCTCTCCTCGTC-----CCCTCCCTTCTCCA	33
HuPSGen-13	GGCACGAGGCTTGAGCGCAGAAACACTTACTTTTCCCCCTACCTGCTCCTCCTCTCCA	60
	*** * * * *	
RatPSGen-13	CTGCAGCCTTTTCTCTTAGCCCCGAACCACTTCCTTCTTCTGCTTGTTCCTCCCTAGGGCGC	93
HuPSGen-13	CAGCCGTCTTTTCTCTTTGCCTCAGCCACTTCCTTCTTCTGCGCTCACCTCCCCAGTGCAC	120
	* * * * *	
RatPSGen-13	GGAGCTGAGTGCAGGGTTCAGACCCACGCGGCGAGCAGCTCTTCAGTGAAGAAGGAAGC	153
HuPSGen-13	TGAAGAAGGTAACCGGGTCCAGACCCACGCGGCGC-CAGTTCTCCGGCGGGAAGGAAAAAC	179
	**** * * * * *	
RatPSGen-13	AAT-CGGAGGGTCAGCAATGAACGTGGAGCATGAGGTAACTCCTGGTGGAGGAAATTC	212
HuPSGen-13	CGCGCAGAGAGGCAGCAATGAATGTGGATCACGAGGTAACTCTTAGTGGAGGAAATTC	239
	* * * * *	
RatPSGen-13	ATCGTCTGGGTTCAAAAATGCCGATGGGAAACTGAGTGTGAAGTTTGGGGTCTCTTCC	272
HuPSGen-13	ATCGTTTGGGTTCAAAAATGCTGATGGAAAGTTAAGCGTGAATTTGGGGTCTCTTCC	299

RatPSGen-13	AAGACGACAGATGTGCCAATCTCTTTGAAGCGTTGGTGGGAACTCTGAAAGCCGCAAAAC	332
HuPSGen-13	GTGATGATAAATGTGCCAACCTCTTTGAAGCATTGGTAGGAACTCTTAAAGCTGCAAAAC	359
	* * * * *	
RatPSGen-13	GAAGGAAGATTGTTACGTACGCAGGAGAGCTGCTTTTGCAAGGTGTTTCATGATGATGTTG	392
HuPSGen-13	GAAGGAAGATTGTAACATATCCAGGAGAGCTGCTTCTGCAAGGTGTTTCATGATGATGTTG	419

RatPSGen-13	ACATTGTATTGCTGCAAGATTAATGTGGTTTGAGATCTGGGGGTA-----	438
HuPSGen-13	ACATTATATTACTGCAAGATTAATGTGGTTTACATATCTTTATGTACTGCCATTTTGT	479

RatPSGen-13	-TCTGGTAAACTGGAATAATTAAGTTAAAGGACAAACAT---GAAGTTCCTTATGTATTT	494
HuPSGen-13	TTCTGGTAAACTGGAATA-TAAAGTGAAAGAACAAACATTTGAACATACTTAATGTATTT	538

RatPSGen-13	TTATAGACCTTTGTAAACAAAAGGGGACTTGT--TGAGAAAGTC---CTGTTTTTATACC	548
HuPSGen-13	TTATAGAAGCTTTGTAAACGAAAGGAGATTTCATGTTTTAGAAAGTCTGTCTTTTTTATATC	598

RatPSGen-13	TTGGAGCAAAACATTACAATGTAAAAATAAAACAAACCTGTTATTTTTTTTTTCTTAAAGA	608
HuPSGen-13	TTGAAAGAAAATCT---ATGTATGATGCTATAAAATAAATCCTATTATTTTTTCTCAGGA	654
	*** * * * *	
RatPSGen-13	AGGTAATCGGAGACGTAGGCAATAAAATGTTTTTCAGAGGTGCGAAAAAGCTTTTGTTTT	668
HuPSGen-13	ATCTGGTTAGGAATTGCAGGCAATGAGATTTTTTGCGGGGCAGGGATGGGAATGTTGTT	714
	* . * * *	
RatPSGen-13	CTTAAACCATTTCT-TAGTCTCTGCC-ACACTTGACACTCCGTCAAAGTGAGAAGCGAACT	726
HuPSGen-13	CATAAATAATTAGACATTTTCTATAGATATTTGACATTCTCGAAAGCAACAAGCAAACT	774
	* * * * *	
RatPSGen-13	AAAGACCAACTGCGGTGGAAAATATTATG---TTTATGTAATAAAAAAATCA-TGT--	780
HuPSGen-13	GAAGACCAACTCCTATGAGAAATATTATGATGTTTATGTAATAAGACATGTAAGTGTCT	834

RatPSGen-13	-	
HuPSGen-13	T 835	

Figure 5

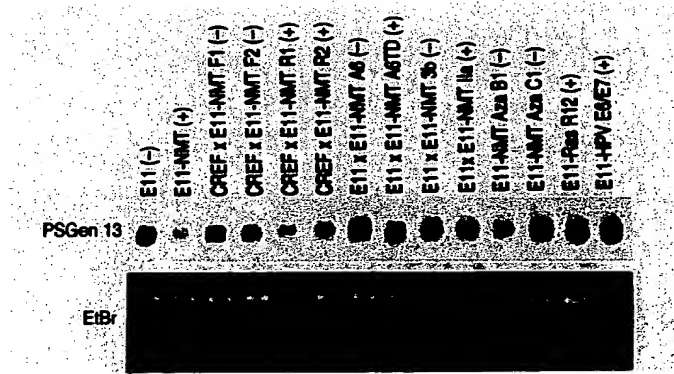


Figure 6

PSGen 13 Suppresses the Transformed Phenotype in E11-NMT Cells

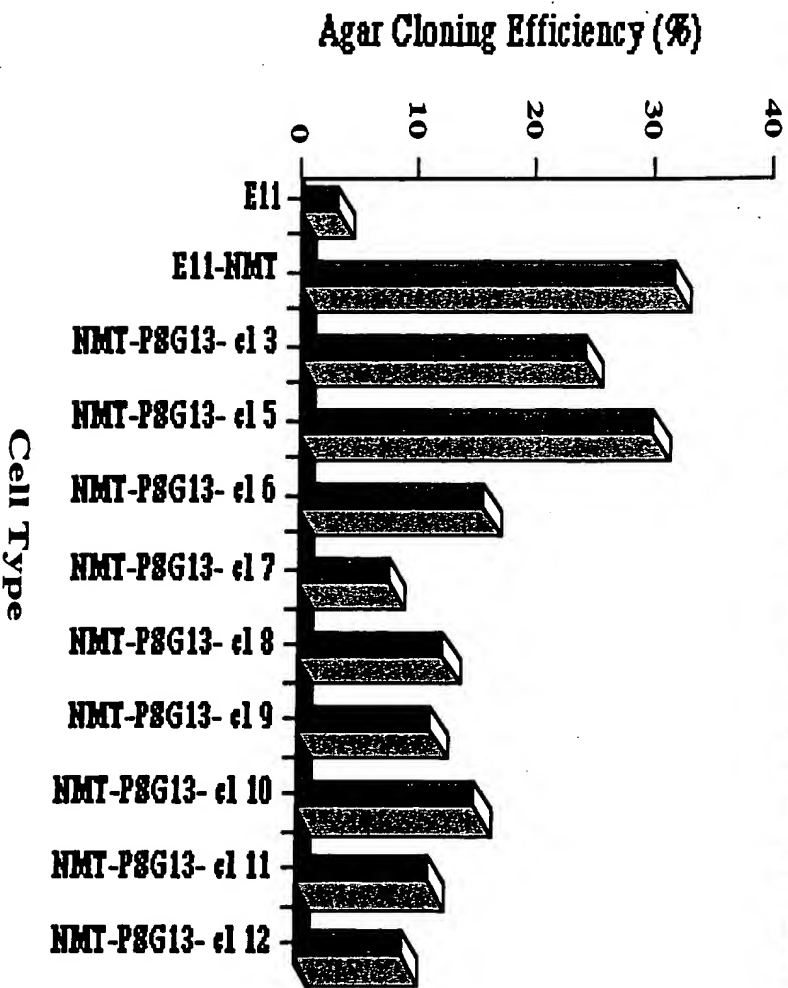


Figure 7

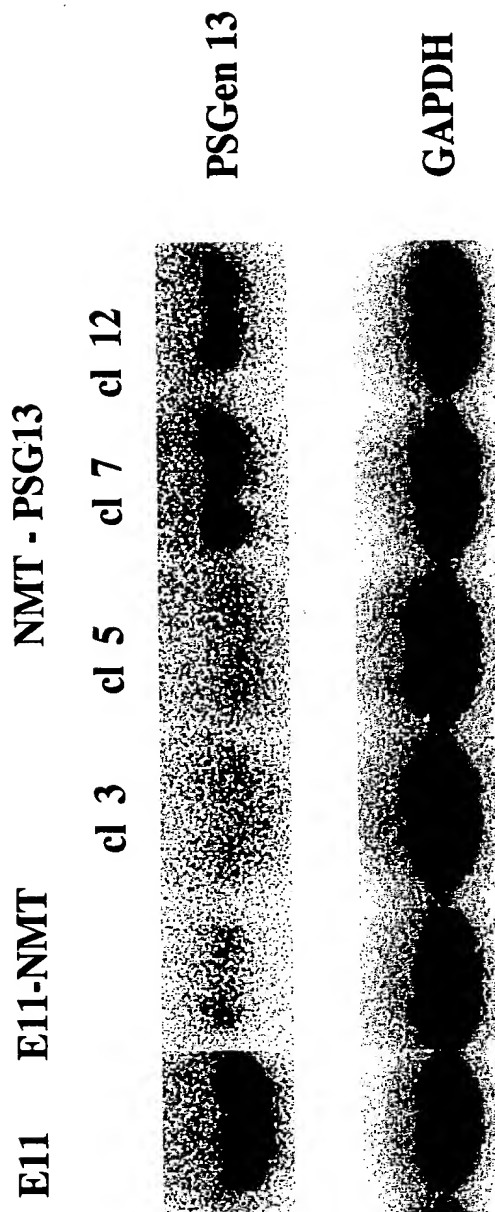
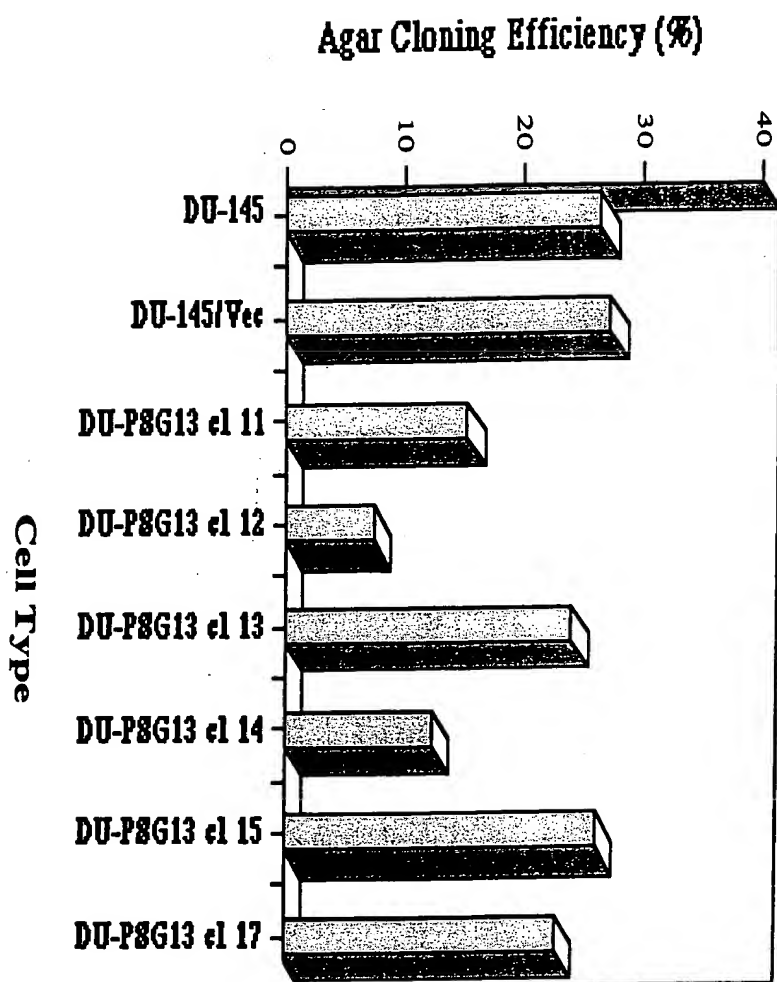


Figure 8

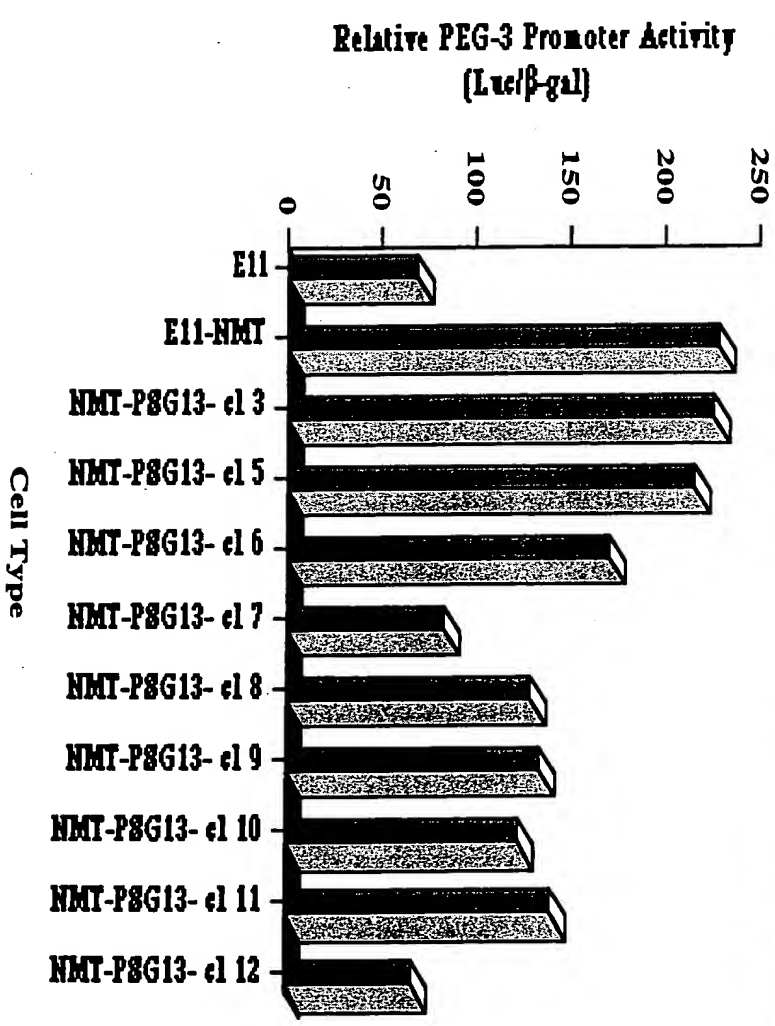
Rat PSGen 13 Inhibits Anchorage Independent Growth in DU-145 Cells



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Figure 9

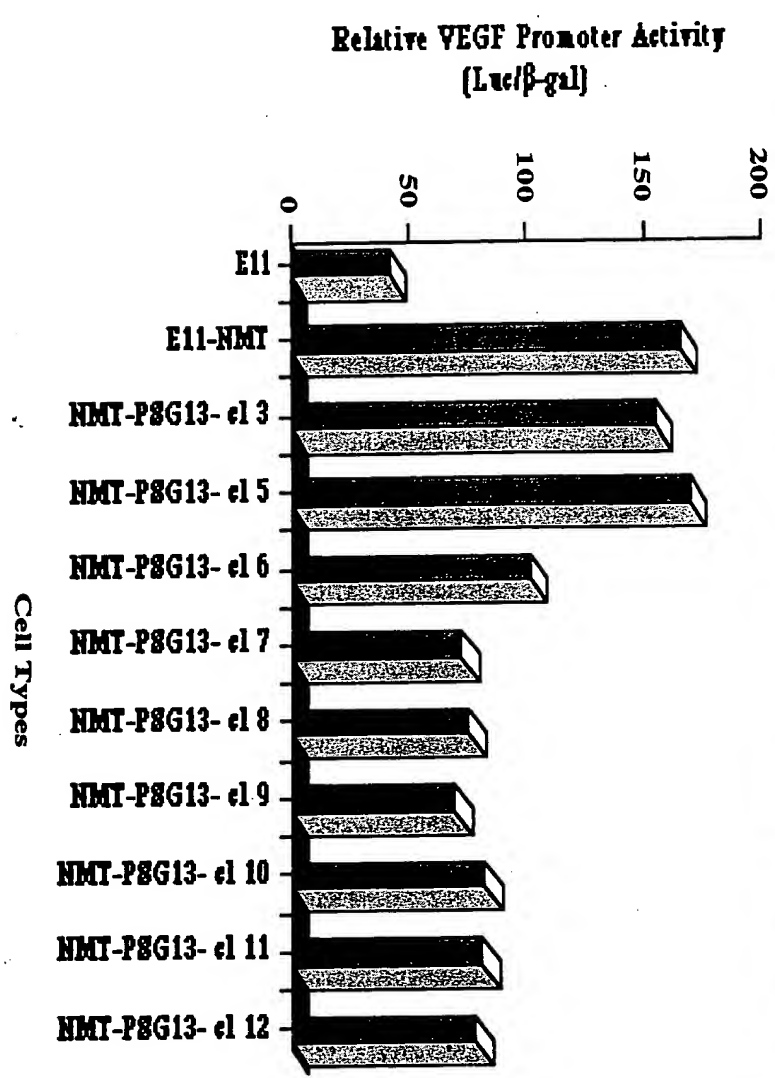
PSGen 13 Suppresses PEG-3 Promoter Activity in E11-NMT Cells



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Figure 10

PSGen 13 Suppresses VEGF Promoter Activity in E11-NMT Cells



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